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TOPOLOGY: US-08-976-259-58 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 4835 base pairs
TYPE: nucleic acid STRANDEDNESS: LENGTH: linear double

BEST AVAILABLE COPY

US-09-848-852A-3 (1-332) x US-08-976-259-58 (1-4835) Percent Similarity:
Best Local Similarity:
Query Match: Score: Pred. No.: Alignment Scores: 0.00929 108.00 31.69% 20.77% 6.09% Conservative: Mismatches: Indels: Matches: Length 4835 59 31 86 108

42 LeuPheGlnGlySerAspLeuGlyMetAlaGluGluAlaGluArgProGlyGluLysAla 61

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                                                                                                                                                  252
                                                                                                                                                                                                               232 ArgArgArgThrIleThrTyrPheAspSerGlnArgThrLeuAsnArgArgCysProLys 251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             162 -----AspGlnValMetAsnMetTyrGlyAspLeuValMetAspThrVal 176
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             LeuAspPheHisginGlyTrpLysglyTyrPheLysMetAsnValAlaArgGlnAsnAsn 286
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                                                                                                                    ----IleAlaLysTyrLeuGlnAlaGluAlaValLysLysAspArg 266
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-AAAGATTATTTCAAG----
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----GCACATGATCGTAAT 1970

Percent Similarity:
Best Local Similarity:
Query Match:

93.00 32.96% 20.95% 5.25% 0.169

Conservative: Mismatches: Indels:

Matches: Length:

Gaps:

score: Pred. No.: Alignment Scores:

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                                         FEATURE:
                                                                                     FEATURE:
                                                                                                                              ORIGINAL SOURCE:
                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 1983 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                           NAME/KEY:
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           LOCATION:
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                                                                                                                                                                                                                                                                                                                            NAME: WEGNER, Harold C.
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1422 TGTTGCAACGGCATTAAATAAGGTGGTGATGGCCGAGCCAGAAAATGACTCC 1473

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                                                                       1362 CGGCGCAGATGCCGCAATTAAGCGCGCTAAAGATGATTACGCTCAAGGTGAATACCGCTT 1421
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                                           276 rPhe---
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                                                                                                                  -----AlaGluAlaValLysLysAspArgLeuAspPheH1sGlnGlyTrpLysGlyTy 276
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---LysMetAsnValAlaArgGlnAsnAsnAspSer 288
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Alignment Scores:
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US-08-752-929-2
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                                                                           FEATURE:
                                                                                                                                                                  MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                               ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                              nucleic acid
EDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                  WEGNER, Harold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20007-5109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08752929
                                                                                                                                                                                                                                        1983 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3: Foley & Lardner
                                                                                                                                                                                                                                                                                                       (202)672-5399
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                                                                                         1..1983
                                             mat_peptide
1..1983
                                                                                                                                       BP-1625
                                                                                                                                                                                                linear
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YAMADA, Akiko
KATO, Seishi
                                                                                                                                           Shewanella putrefaciens SCRC-2874 (FERM
                                                                                                                                                                             DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-JAN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene Coding for Eicosapentaenoic Acid Synthesizing
Enzymes and Process for Production of Eiscosapentaenoic
Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                              JP 4-147945
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Query Match:

Percent Similarity: Best Local Similarity:

0.169 93.00 32.96% 20.95% 5.25%

Length:
Matches:
Conservative:
Mismatches:
Indels:

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